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European Patent Office Erhardtstrasse 27 D-80298 München Germany

For the attention of International Preliminary Examining Authority

Our Ref:

P33779WO/TF/L.IP

Your Ref:

22 June 2004

Dear Sirs

International (PCT) Patent Application No. PCT/GB2003/005158 In the name of Medical Research Council et al

I file herewith a Demand for Chapter II PCT.

Arrangements are being made separately by our formalities department for payment of the associated fees. In the event of non-payment or underpayment please debit the requisite sum from our deposit account number 28050204.

An amended set of claims is enclosed, in which claims 1 and 2 have been amended. Claim 1 has been amended to replace "A crystal structure of the pRb/E2F₍₄₀₉₋₄₂₆₎ complex," with "A crystal comprising a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, wherein the crystal structure is". Claim 2 has been amended to refer to "A crystal as claimed in claim 1" in view of the amendment made to claim 1.

These amendments have been made to claims 1 and 2 of the present application in light of the International Searching Authority being of the opinion that the subject matter of previous claims 1 and 2 related to the presentation of information. I submit that the subject matter of amended claims 1 and 2 clearly does not relate to the presentation of information because these claims now define a crystal per se.

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Kilburn & Strode

Yours faithfully

FORD, Timothy James Agent for the Applicants

Enc.

Chapter II Demand

EPO Form 1037

Copy letter to International Bureau

Copy claims filed at International Bureau



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JC20 Rec'd PCT/P1 2 7 MAY 2005

Claims

- 1. A crystal comprising a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, wherein the crystal structure is characterised by the atomic co-ordinates of Annex 1.
- 2. A crystal as claimed in claim 1, wherein the interactions between E2F₍₄₀₉₋₄₂₆₎ and pRb comprise one or more of the following interactions:

E2F ₍₄₀₉₋₄₂₆₎ residue	pRb residue
Leu ₄₀₉	Lys ₅₄₈
Tyr ₄₁₁	Glu ₅₅₁
Tyr ₄₁₁	Ile ₅₃₂
Tyr ₄₁₁	Glu ₅₅₄
His ₄₁₂	Arg ₆₅₆
His ₄₁₂	Lys ₆₅₃
Gly ₄₁₄	Glu ₅₃₃
Gly ₄₁₄	Lys ₆₅₂
Leu ₄₁₅	Leu ₆₄₉
Leu ₄₁₅	Glu ₅₅₃
Leu ₄₁₅	Lys ₅₃₇
Glu ₄₁₇	Lys ₅₃₇
Gly ₄₁₈	Arg ₄₆₇
Glu ₄₁₉	Thr ₆₄₅
Arg ₄₂₂	Glu ₄₆₄
Asp ₄₂₃	Arg ₄₆₇
Leu ₄₂₄	Lys ₅₃₀
Phe ₄₂₅	Phe ₄₈₂
Phe ₄₂₅	Lys ₄₇₅

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- 3. A method to identify an agent which modulates the interaction between pRb and E2F₍₄₀₉₋₄₂₆₎, the method comprising:
- a) combining together pRb, E2F₍₄₀₉₋₄₂₆₎ and an agent, under conditions in which pRb and E2F₍₄₀₉₋₄₂₆₎ form a complex;
 - b) growing a crystal structure of any pRb/ E2F₍₄₀₉₋₄₂₆₎ complex; and
- c) analysing the crystal to determine whether the agent is an agent which modulates
 the interaction between pRb and E2F₍₄₀₉₋₄₂₆₎.
 - 4. A method, as claimed in claim 3, wherein the combining of the components is pRb with the agent and then E2F₍₄₀₉₋₄₂₆₎.
- A method as claimed in claim 3, wherein the combining of the components is
 E2F₍₄₀₉₋₄₂₆₎ with the agent and then pRb.
 - 6. A method as claimed in claim 3, wherein the combining of the components is pRb with E2F₍₄₀₉₋₄₂₆₎ and then the agent.
 - 7. A method as claimed in any one of claims 3 to 6, wherein step c) comprises comparing the crystal structure to the crystal structure of claim 1
- 8. A method as claimed in any one of claims 3 to 7, wherein the agent is selected using the three dimensional atomic co-ordinates of Annex 1.
 - 9. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising selecting an agent using the three-dimensional atomic coordinates of Annex 1.

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- 10. A method as claimed in claim 9, wherein said selection is performed in conjunction with computer modeling.
 - 11. A method as claimed in claim 9 or 10, wherein the method further comprises the steps of:
 - a) contacting the selected agent with pRb and E2F₍₄₀₉₋₄₂₆₎ under conditions in which pRb and E2F₍₄₀₉₋₄₂₆₎ can form a complex; and
 - b) measuring the binding affinity of pRb to E2F₍₄₀₉₋₄₂₆₎ in the presence of the agent and comparing the binding affinity to that of pRb to E2F₍₄₀₉₋₄₂₆₎ when in the absence of the agent, wherein an agent modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex when there is a change in the binding affinity of pRb to E2F₍₄₀₉₋₄₂₆₎ when in the presence of the agent.
 - 12. A method as claimed in claim 11, wherein the method further comprising:
- a) growing a supplementary crystal from a solution containing pRb and E2F₍₄₀₉₋₄₂₆₎ and the selected agent where said agent changes the binding affinity of the pRb/E2F₍₄₀₉₋₄₂₆₎ complex under conditions in which pRb and E2F₍₄₀₉₋₄₂₆₎ can form a complex;
 - b) determining the three-dimensional atomic coordinates of the supplementary crystal by X-ray diffraction using molecular replacement analysis;

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- c) comparing the three dimensional coordinates with those for the crystal structure as claimed in claim 1; and
- d) selecting a second generation agent using the three-dimensional atomic coordinates determined for the supplementary crystal.
- 13. A method as claimed in claim 12, wherein said selection is performed in conjunction with computer modeling.
- 14. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising:

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- a) contacting a selected agent with pRb and E2F₍₄₀₉₋₄₂₆₎ under conditions in which pRb and E2F₍₄₀₉₋₄₂₆₎ can form a complex; and
- b) measuring the binding affinity of pRb to E2F₍₄₀₉₋₄₂₆₎ in the presence of the agent and comparing the binding affinity to that of pRb to E2F₍₄₀₉₋₄₂₆₎ when in the absence of the agent, wherein an agent modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex when there is a change in the binding affinity of pRb to E2F₍₄₀₉₋₄₂₆₎ when in the presence of the agent.
- 15. A method as claimed in claim 14, wherein the method further comprising:
- a) growing a supplementary crystal from a solution containing pRb and E2F₍₄₀₉₋₄₂₆₎ and the selected agent where said agent changes the binding affinity of the pRb/E2F₍₄₀₉₋₄₂₆₎ complex under conditions in which pRb and E2F₍₄₀₉₋₄₂₆₎ can form a complex;
 - b) determining the three-dimensional atomic coordinates of the supplementary crystal by X-ray diffraction using molecular replacement analysis;
 - c) comparing the three dimensional coordinates with those for the crystal structure claimed in claim 1; and
 - d) selecting a second generation agent using the three-dimensional atomic coordinates determined for the supplementary crystal.
 - 16. A method as claimed in claim 15, wherein said selection is performed in conjunction with computer modeling.
 - 17. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising:
 - a) selecting an agent;

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- b) co-crystalising pRb with the agent;
- c) determining the three dimensional coordinates of the pRb-agent association by X-ray diffraction using molecular replacement analysis; and





- d) comparing the three dimensional coordinates with those of the crystal structure claimed in claim 1.
- 18. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising:
- a) selecting an agent;
- b) crystalising pRb and soaking the agent into the crystal;
- c) determining the three dimensional coordinates of the pRb-agent association by X-ray diffraction using molecular replacement analysis; and
- d) comparing the three dimensional coordinates with those of the crystal structure claimed in claim 1.
 - 19. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising:
- 15 a) selecting an agent;
 - b) co-crystalising pRb, E2F₍₄₀₉₋₄₂₆₎ and the agent;
 - c) determining the three dimensional coordinates of the pRb-E2F-agent association by X-ray diffraction using molecular replacement analysis; and
 - d) comparing the three dimensional coordinates with those of the crystal structure claimed in claim 1.
 - 20. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising:
 - a) selecting an agent;
- b) co-crystalising pRb and E2F₍₄₀₉₋₄₂₆₎ and soaking the agent into the crystal;
 - c) determining the three dimensional coordinates of the pRb-E2F-agent association by X-ray diffraction using molecular replacement analysis; and
 - d) comparing the three dimensional coordinates with those of the crystal structure claimed in claim 1.



- 21. A method as claimed in any one of claims 17 to 20, wherein the agent is selected using the three dimensional atomic co-ordinates of Annex 1
- 22. A method as claimed in any one of claims 17 to 21, wherein the methods further comprise selecting a second generation agent using the three dimensional atomic coordinates determined in step c).
 - 23. A method as claimed in claim 22, wherein the second generation agent is selected using the three dimensional atomic coordinates of Annex 1.
 - 24. A method as claimed in claim 22 or 23, wherein the selection is performed in conjunction with computer modeling.
- 25. A method of identifying an agent as claimed in any one of claims 3 to 24, wherein the selected agent and/or the second generation agent mimics a structural feature of E2F₍₄₀₉₋₄₂₆₎, when said E2F₍₄₀₉₋₄₂₆₎ is bound to pRb.
 - 26. A method as claimed in claim 9 or 10, wherein method comprises the further steps of:
- a) contacting the selected agent with a pRb/E2F₍₄₀₉₋₄₂₆₎ complex; and
 - b) determining whether the agent affects the stability of the complex.
 - 27. A method as claimed in claim 26, wherein the determination is with fluorescence polarization.
 - 28. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising:
 - a) contacting a fluorescently tagged E2F₍₄₀₉₋₄₂₆₎ peptide (E2F-fluoropeptide) with pRb to allow pRb/E2F-fluoropeptide complex formation;
- 30 b) detecting the fluorescence polarization;



- c) adding a selected agent; and
- d) detecting the fluorescence polarization in the presence of the agent.
- 29. A method of identifying an agent that modulates a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, comprising;
- a) contacting a fluorescently tagged E2F₍₄₀₉₋₄₂₆₎ peptide (E2F-fluoropeptide) with pRb to allow pRb/E2F-fluoropeptide complex formation;
- b) detecting the fluorescence polarization;
- c) contacting a selected agent with pRb and E2F₍₄₀₉₋₄₂₆₎ peptide (E2F-fluoropeptide) under conditions in which pRb and E2F-fluoropeptide can form a complex;
 - d) detecting the fluorescence polarization; and
 - e) comparing the fluorescence polarization detected in b) and d).
- 30. A method as claimed in claim 28 or 29, wherein the fluorescently tagged E2F peptide is selected using the three dimensional atomic co-ordinates of Annex 1.
 - 31. A method as claimed in any one of claims 28 to 30, wherein a decrease in fluorescence polarization in the presence of the agent indicates that the agent destabilises the complex.

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- 32. A method as claimed in any one of claims 28 to 31, wherein the method comprises the further step of adding untagged E2F₍₄₀₉₋₄₂₆₎ and detecting fluorescence polarization.
- 25 33. A method as claimed in claim 32, wherein if fluorescence polarization decreases, on addition of the untagged E2F₍₄₀₉₋₄₂₆₎, the agent does not stabilise the complex.
 - 34. A method as claimed in claim 32 or 33, wherein if there is no substantial change in fluorescence polarization, on addition of the untagged E2F₍₄₀₉₋₄₂₆₎, the agent stabilises the complex.





- 35. A method as claimed in any one of claims 28 to 34, wherein the method further comprises:
- a) contacting a fluorescently tagged E7 peptide (E7-fluoropeptide) with pRb to allow pRb/E7-fluoropeptide complex formation;
- b) detecting the fluorescence polarization;
- c) adding an agent that modulates pRb/E2F(409-426) complex; and
- d) detecting the fluorescence polarization in the presence of the agent.
- 36. A method as claimed in any one of claims 28 to 34, wherein the method further comprises:
 - a) contacting a fluorescently tagged E7 peptide (E7-fluoropeptide) with pRb to allow pRb/E7-fluoropeptide complex formation;
 - b) detecting the fluorescence polarization;
- 15 c) contacting an agent that modulates pRb/E2F₍₄₀₉₋₄₂₆₎ complex with pRb and E7-fluoropeptide under conditions in which pRb and E7-fluoropeptide can from a complex;
 - d) detecting the fluorescence polarization; and
 - e) comparing the fluorescence polarization detected in b) and d).

- 37. A method as claimed in claim 35 or 36, wherein a decrease in fluorescence polarization indicates that the agent also inhibits E7 binding to pRb.
- 38. A method as claimed in any one of claims 11 to 16, wherein the binding affinity is
 measured by isothermal titration calorimetry.
 - 39. A method as claimed in any one of claims 11 to 16, wherein the binding affinity is measure by Surface Plasmon Resonance (SPR).



- 40. An agent, that modulates the interaction between pRb and E2F₍₄₀₉₋₄₂₆₎, identified by a method as claimed in any one of claims 3 to 39.
- 41. An agent, as claimed in claim 40, for use as an apoptosis promoting factor in the prevention or treatment of proliferative diseases.
 - 42. An agent as claimed in claim 40 or 41, wherein the agent is for use in preventing or treating cancer, which may be pancreatic cancer and related diseases.
- 43. The use of an agent, which modulates the formation of a pRb/E2F₍₄₀₉₋₄₂₆₎ complex, identified by a method as claimed in any one of claims 3 to 39, in the manufacture of a medicament for the prevention or treatment of proliferative diseases.
- 44. The use of an agent as claimed in claim 43, wherein the proliferative diseases are cancer, preferably pancreatic cancer and related diseases.
 - 45. The use of the atomic co-ordinates of the crystal structure as claimed in claim 1 or 2, for identifying an agent that modulates the formation of a pRb/E2F₍₄₀₉₋₄₂₆₎ complex.

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46. Computer readable media comprising a data storage material encoded with computer readable data, wherein said computer readable data comprises a set of atomic co-ordinates of the pRb/E2F₍₄₀₉₋₄₂₆₎ crystal structure of Annex 1 recorded thereon.